0φcD 04-10-01

②

RAW SEQUENCE LISTING   DATE: 03/30/2001
PATENT APPLICATION: US/09/804,472   TIME: 11:17:56

Input Set : A:\Seqlist.txt
Output Set: N:\CRF3\03302001\I804472.raw

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   4 <110> APPLICANT: SHAO, Wei et al.
   6 <120> TITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS,
   7       NUCLEIC ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS,
   8       AND USES THEREOF
  10 <130> FILE REFERENCE: CL001163
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C--> 12 <141> CURRENT FILING DATE: 2001-03-13
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  14 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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  18 <212> TYPE: DNA
  19 <213> ORGANISM: Human
  21 <400> SEQUENCE: 1
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  24 tcttttttctt tttcttcttc tatttaaaaa ttctaatcat ggatgcttct tccgaccctt 180
  25 atttgcctta tgacggggga ggagacaata ttccctgag ggaattacat aaaagaggaa 240
  26 ctcattatac aatgacaaat ggaggcagca ttaacagttc tacacattta ctggatcttt 300
  27 tggatgaacc aattccaggt gttggtacat atgatgattt ccatactatt gattgggtgc 360
  28 gagaaaaatg taaagacaga gaaaggcata gacggatcaa cagcaaaaag aaagaatcag 420
  29 catgggaaat gacaaaaagt ttgtatgatg cgtggtcagg atggctagta gtaacactaa 480
  30 caggattggc atcaggggca ctggccggat taatagacat tgctgccgat tggatgactg 540
  31 acctaaagga gggcatttgc cttagtgcgt tgtggtacaa ccacgaacag tgctgttggg 600
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  33 aattaatcat aggtcaagca gagggtcctg gttcttatat catgaactac ataatgtaca 720
  34 tcttctgggc cttgagtttt gcctttcttg cagtttccct ggtaaaggta tttgctccat 780
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  36 gttacttggg aaaatggact ttaatgatta aaaccatcac attagtcctg gctgtggcat 900
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  38 tcttttccta cctctttcca aagtatagca caaacgaagc taaaaaaagg gaggtgctat 1020
  39 cagctgcctc agctgcaggg gtttctgtag cttttggtgc accaattgga ggagttcttt 1080
  40 ttagcctgga agaggttagc tattattttc ctctcaaaac tttatggaga tcattttttg 1140
  41 ctgctttagt ggctgcattt gtttttgaggt ccatcaatcc atttggtaac agccgtctgg 1200
  42 tcctttttta tgtggagtat catacaccat ggtacctttt tgaactgttt cctttttattc 1260
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  47 gaaatgacat gaatgccagt aaaattgtcg atgacattcc tgatcgtcca gcaggcattg 1560
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  50 cgatcgcagg aaggattgtg gggattgcgg tggagcagct tgcctactat caccacgact 1740
  51 ggtttatctt taaggagtgg tgtgaggtcg gggctgattg cattcacct ggcctttatg 1800
  52 ccatggttgg tgctgctgca tgcttaggtg gtgtgacaag aatgactgtc tccctggtgg 1860
  53 ttattgtttt tgagcttact ggaggcttgg aatatattgt tccccttatg gctgcagtca 1920
  54 tgaccagtaa atgggttgga gatgcctttg gcagggaagg catttatgaa gcacacatcc 1980
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RAW SEQUENCE LISTING   DATE: 03/30/2001
PATENT APPLICATION: **US/09/804,472**   TIME: 11:17:56

Input Set : **A:\Seqlist.txt**
Output Set: **N:\CRF3\03302001\I804472.raw**

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63 taacaaaaaa agatatcctc cggcatatgg cccagacggc aaaccaagac cccgcttcaa 2520
64 taatgttcaa ctgaatctca cagatgagga gagagaagaa acggaagagg aagtttattt 2580
65 gttgaatagc acaactcttt aacctgaggg agtcatctac ttttttttcc tcctttacaa 2640
66 aaaaagaaag gaaatataaa agccgggttt ttgcaacatg gtttgcaaat aatgctggtg 2700
67 gaatggagga gttgtttggg gagggaaagg agagagaagg aaaggagtga ggtatttccc 2760
68 gtctaacaga aagcagcgta tcaactccta ttgttctgca ctggatgcat tcagctgagg 2820
69 atgtgcctga tagtgcaggc ttgcgcctca acagagatga cagcagagtc ctcgagcacc 2880
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71 tttgaattga gccatctata aaactgcaag gtcttgccct ttttttttaat caaaactgtt 3000
72 ctgtttaatt catgaattgt atagttaagc attacctttc tacattccag aagagccttt 3060
73 atttctctct ctctctctct ctctctctct ctctctactg agctgtaaca aagcctcttt 3120
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78 tgaagcttgg ttttaaagga taaagttttc tttttgttt tcctctcaga ctttatggat 3420
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87 <213> ORGANISM: Human
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92 Asn Ile Pro Leu Arg Glu Leu His Lys Arg Gly Thr His Tyr Thr Met
93              20                  25                  30
94 Thr Asn Gly Gly Ser Ile Asn Ser Ser Thr His Leu Leu Asp Leu Leu
95          35                  40                  45
96 Asp Glu Pro Ile Pro Gly Val Gly Thr Tyr Asp Asp Phe His Thr Ile
97      50                  55                  60
98 Asp Trp Val Arg Glu Lys Cys Lys Asp Arg Glu Arg His Arg Arg Ile
99 65                  70                  75                  80
100 Asn Ser Lys Lys Lys Glu Ser Ala Trp Glu Met Thr Lys Ser Leu Tyr
101              85                  90                  95
102 Asp Ala Trp Ser Gly Trp Leu Val Val Thr Leu Thr Gly Leu Ala Ser
103              100                 105                 110
104 Gly Ala Leu Ala Gly Leu Ile Asp Ile Ala Ala Asp Trp Met Thr Asp
105          115                 120                 125
```

```
106 Leu Lys Glu Gly Ile Cys Leu Ser Ala Leu Trp Tyr Asn His Glu Gln
107     130             135             140
108 Cys Cys Trp Gly Ser Asn Glu Thr Thr Phe Glu Glu Arg Asp Lys Cys
109 145             150             155             160
110 Pro Gln Trp Lys Thr Trp Ala Glu Leu Ile Ile Gly Gln Ala Glu Gly
111             165             170             175
112 Pro Gly Ser Tyr Ile Met Asn Tyr Ile Met Tyr Ile Phe Trp Ala Leu
113             180             185             190
114 Ser Phe Ala Phe Leu Ala Val Ser Leu Val Lys Val Phe Ala Pro Tyr
115         195             200             205
116 Ala Cys Gly Ser Gly Ile Pro Glu Ile Lys Thr Ile Leu Ser Gly Phe
117     210             215             220
118 Ile Ile Arg Gly Tyr Leu Gly Lys Trp Thr Leu Met Ile Lys Thr Ile
119 225             230             235             240
120 Thr Leu Val Leu Ala Val Ala Ser Gly Leu Ser Leu Gly Lys Glu Gly
121             245             250             255
122 Pro Leu Val His Val Ala Cys Cys Cys Gly Asn Ile Phe Ser Tyr Leu
123             260             265             270
124 Phe Pro Lys Tyr Ser Thr Asn Glu Ala Lys Lys Arg Glu Val Leu Ser
125         275             280             285
126 Ala Ala Ser Ala Ala Gly Val Ser Val Ala Phe Gly Ala Pro Ile Gly
127     290             295             300
128 Gly Val Leu Phe Ser Leu Glu Glu Val Ser Tyr Tyr Phe Pro Leu Lys
129 305             310             315             320
130 Thr Leu Trp Arg Ser Phe Phe Ala Ala Leu Val Ala Ala Phe Val Leu
131             325             330             335
132 Arg Ser Ile Asn Pro Phe Gly Asn Ser Arg Leu Val Leu Phe Tyr Val
133             340             345             350
134 Glu Tyr His Thr Pro Trp Tyr Leu Phe Glu Leu Phe Pro Phe Ile Leu
135         355             360             365
136 Leu Gly Val Phe Gly Gly Leu Trp Gly Ala Phe Phe Ile Arg Ala Asn
137     370             375             380
138 Ile Ala Trp Cys Arg Arg Arg Lys Ser Thr Lys Phe Gly Lys Tyr Pro
139 385             390             395             400
140 Val Leu Glu Val Ile Ile Val Ala Ala Ile Thr Ala Val Ile Ala Phe
141             405             410             415
142 Pro Asn Pro Tyr Thr Arg Leu Asn Thr Ser Glu Leu Ile Lys Glu Leu
143             420             425             430
144 Phe Thr Asp Cys Gly Pro Leu Glu Ser Ser Ser Leu Cys Asp Tyr Arg
145         435             440             445
146 Asn Asp Met Asn Ala Ser Lys Ile Val Asp Asp Ile Pro Asp Arg Pro
147     450             455             460
148 Ala Gly Ile Gly Val Tyr Ser Ala Ile Trp Gln Leu Cys Leu Ala Leu
149 465             470             475             480
150 Ile Phe Lys Ile Ile Met Thr Val Phe Thr Phe Gly Ile Lys Val Pro
151             485             490             495
152 Ser Gly Leu Phe Ile Pro Ser Met Ala Ile Gly Ala Ile Ala Gly Arg
153             500             505             510
154 Ile Val Gly Ile Ala Val Glu Gln Leu Ala Tyr Tyr His His Asp Trp
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156 Phe Ile Phe Lys Glu Trp Cys Glu Val Gly Ala Asp Cys Ile Thr Pro
157      530                    535                    540
158 Gly Leu Tyr Ala Met Val Gly Ala Ala Ala Cys Leu Gly Gly Val Thr
159 545                    550                    555                    560
160 Arg Met Thr Val Ser Leu Val Val Ile Val Phe Glu Leu Thr Gly Gly
161                    565                    570                    575
162 Leu Glu Tyr Ile Val Pro Leu Met Ala Ala Val Met Thr Ser Lys Trp
163                    580                    585                    590
164 Val Gly Asp Ala Phe Gly Arg Glu Gly Ile Tyr Glu Ala His Ile Arg
165              595                    600                    605
166 Leu Asn Gly Tyr Pro Phe Leu Asp Ala Lys Glu Glu Phe Thr His Thr
167      610                    615                    620
168 Thr Leu Ala Ala Asp Val Met Arg Pro Arg Arg Asn Asp Pro Pro Leu
169 625                    630                    635                    640
170 Ala Val Leu Thr Gln Asp Asn Met Thr Val Asp Asp Ile Glu Asn Met
171                    645                    650                    655
172 Ile Asn Glu Thr Ser Tyr Asn Gly Phe Pro Val Ile Met Ser Lys Glu
173              660                    665                    670
174 Ser Gln Arg Leu Val Gly Phe Ala Leu Arg Arg Asp Leu Thr Ile Ala
175         675                    680                    685
176 Ile Glu Ser Ala Arg Lys Lys Gln Glu Gly Ile Val Gly Ser Ser Arg
177      690                    695                    700
178 Val Cys Phe Ala Gln His Thr Pro Ser Leu Pro Ala Glu Ser Pro Arg
179 705                    710                    715                    720
180 Pro Leu Lys Leu Arg Ser Ile Leu Asp Met Ser Pro Phe Thr Val Thr
181                    725                    730                    735
182 Asp His Thr Pro Met Glu Ile Val Val Asp Ile Phe Arg Lys Leu Gly
183              740                    745                    750
184 Leu Arg Gln Cys Leu Val Thr His Asn Gly Arg Leu Leu Gly Ile Ile
185         755                    760                    765
186 Thr Lys Lys Asp Ile Leu Arg His Met Ala Gln Thr Ala Asn Gln Asp
187      770                    775                    780
188 Pro Ala Ser Ile Met Phe Asn
189 785                    790
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205 tttaccggga ggatttcccc catcagtgag tgctgactgt cattttcatt ctttatgatc 180
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RAW SEQUENCE LISTING   DATE: 03/30/2001
PATENT APPLICATION: US/09/804,472   TIME: 11:17:56

Input Set : A:\Seqlist.txt
Output Set: N:\CRF3\03302001\I804472.raw

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216 gccctctgag ttttctcttt ccttctctca cctccaattt tcatgtaaac actttcagtt 840
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FYI.

VERIFICATION SUMMARY   DATE: 03/30/2001
PATENT APPLICATION: US/09/804,472   TIME: 11:17:57

Input Set : A:\Seqlist.txt
Output Set: N:\CRF3\03302001\I804472.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
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L:285 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
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```
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L:681 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:682 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
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